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2 **Interpretive summary.** Use of on-farm data to guide treatment and control of mastitis caused by  
3 *Streptococcus uberis*, by Samson *et al.* To reduce the risk of antimicrobial resistance, judicious use of  
4 antimicrobials is advocated. We show that routinely available DHI and treatment data can be used in  
5 veterinary practice to predict cure of *S. uberis* mastitis. Probability of apparent cure is higher among  
6 1st and 2nd parity animals compared to older cows, and in animals with short-duration elevated SCC  
7 compared to those with repeated SCC elevation before occurrence of mastitis. This knowledge  
8 enables farmers and veterinarians to tailor antimicrobial use for treatment of mastitis, and to put  
9 increased emphasis on prevention of cases with poor prognosis.

10

11 **Use of on-farm data to guide treatment and control of mastitis caused by *Streptococcus uberis***

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### ABSTRACT

Treatment of mastitis is the most common reason for use of antimicrobials in dairy cattle. The responsible use of antimicrobials could be strengthened by knowledge of predictors for cure, which would help to tailor treatment decisions. Ideally, to allow for widespread uptake, this would be achieved using data that are routinely available. To assess whether this is feasible in practice, farmers were invited to submit milk samples from mastitis cases to their veterinary practice for bacteriological culture. Among 624 culture-positive samples, 251 were positive for *Streptococcus uberis*. Using cow-level data, cases were classified as severe clinical mastitis (CM; “severe”), 1<sup>st</sup> non-severe CM (“first”), repeated non-severe CM (“repeat”), or subclinical mastitis (“subclinical”). Additional data were collected at cow-level (somatic cell count (SCC), parity, lactation stage, milk yield, fat and protein content, treatment) and at herd-level (housing, bedding, pre-milking teat disinfection, post-milking teat disinfection). Severe cases were overrepresented among heifers and animals in early lactation whereas repeat cases were overrepresented in cows with 3 or more lactations. The probability of cure was higher among 1<sup>st</sup> and 2<sup>nd</sup> parity animals than among older cows, and higher in animals with a single elevated cow-level SCC than in animals with multiple high SCC records. Results obtained in the current study are similar to those previously described for *Staphylococcus aureus* mastitis. Thus, routinely available cow-level information can help to predict the outcome of antimicrobial treatment of the most common causes of Gram-positive mastitis.

53 **Key words:** mastitis, *Streptococcus uberis*, prognosis, antimicrobial treatment, clinical manifestation

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## INTRODUCTION

Antimicrobial resistance is increasingly perceived as a threat to human and animal health and the need for responsible use of antimicrobials is emphasized by a range of national and international bodies (UK Department of Health 2013; World Health Organisation 2015). Key elements of the approach proposed by the World Health Organisation (WHO) include reduction of the incidence of infection and optimized use of antimicrobial medicines (WHO, 2015). Farmers are increasingly aware of the need to use antimicrobials responsibly. In a recent survey in the UK over 70% of dairy farmers said that reducing antibiotic usage would be “a good thing to do” (Jones *et al.*, 2015). Veterinarians can play an important role in this process by providing information on ways to achieve reductions in antibiotic usage, e.g. by minimizing the risk of disease or through development of treatment protocols (Raymond *et al.*, 2006; Jones *et al.*, 2015).

On dairy farms, treatment of mastitis is a major reason for use of antimicrobials. For example, Pol and Ruegg (2007) calculated the estimated overall exposure to antimicrobial drugs of cattle on conventional dairy farms as 5.43 defined daily doses (DDD) per cow per year. This included 3.58 DDD of intramammary applications (2.02 DDD during lactation and 1.56 DDD at dry off) and 1.85 DDD of parenteral use. Clinical mastitis (CM) was the most common reason for intramammary or parenteral antimicrobial usage. To reduce the use of antimicrobials protocols for selective treatment of dry cows and cattle with CM have been developed, including protocols based on culture and on cow-factors such as SCC (Lago *et al.*, 2011; Cameron *et al.*, 2014; Scherpenzeel *et al.*, 2014). In culture-based protocols, treatment decisions are largely based on the distinction between gram-positive growth, gram-negative growth and no growth (Lago *et al.*, 2011; Cameron *et al.*, 2014). Further refinement of treatment decisions may be possible when pathogen factors, such as antimicrobial resistance, and host characteristics, including duration of infection and parity, are taken into account, but this has only been described in detail for *Staphylococcus aureus* (Barkema *et al.*, 2006).

80           In many countries on different continents *Streptococcus uberis* is among the most common  
81 gram-positive causes of CM (Olde Riekerink *et al.*, 2008; Petrovski *et al.*, 2011; Verbeke *et al.*, 2014).  
82 The organism is also responsible for a considerable proportion of subclinical mastitis cases (Bradley  
83 *et al.*, 2007; Sampimon *et al.*, 2009). Intramammary infections and CM caused by *S. uberis* can be  
84 transient, recurrent or chronic and a wide range of cure rates has been reported in response to  
85 treatment (Zadoks *et al.*, 2003; Zadoks 2007). Despite its importance as a mastitis pathogen little is  
86 known about risk factors for the clinical manifestation or treatment outcome of *S. uberis* IMI.  
87 Therefore, the aim of this study was to generate data that could inform guidelines for improved  
88 management of *S. uberis* mastitis under field conditions. To that end, we conducted a farm-based  
89 study of herd- and cow-level risk factors that are associated with the clinical manifestation and  
90 likelihood of apparent cure of *S. uberis* IMI as based on SCC. In doing so, we only used tools and data  
91 that are routinely available to farmers and veterinary practices, including treatment and DHI records,  
92 because our aim was to generate low-cost guidelines for improved management of *S. uberis* mastitis  
93 under field conditions.

94

**MATERIALS AND METHODS**

95

***Milk sampling and Bacteriological culture***

97 From August 2012 until January 2014, quarter milk samples (n=624) were collected from French  
98 dairy cows with clinical or subclinical mastitis using standard aseptic sample collection methods  
99 (National Mastitis Council, 1999). Detection and sampling of mastitis cases was driven by  
100 participating farmers. To motivate farmers to participate in the study, all clients of our veterinary  
101 practice (Vetformance, Villaines la Juhel, France) with more than 50 lactating cows (approximately  
102 500 farms), received an invitation to sample clinical and subclinical mastitis cases at their farms. The  
103 bacteriological analysis of the samples was free of charge for the farmers. In addition, upon return of  
104 completed data information sheets, a head collar for a cow was offered to the farmers. One hundred  
105 and forty two farmers submitted at least one milk sample, indicating an uptake of approximately  
106 28%.

107 Milk samples were subjected to bacteriological culture in the laboratory of Vetformance.  
108 Aliquots of milk (10 µl) were plated onto three media, i.e. (1) Colombia blood agar containing 5%  
109 sheep blood (bioMérieux, Craponne, France; Ref. 43041); (2) Colistin Nalidixic Acid (CNA) agar (blood  
110 agar plate containing 5% sheep blood, Colistin (10 mg/L) and Nalidixic Acid (15 mg/L) (bioMérieux;  
111 Ref. 43071) and (3) Bromo Cresol Purple (BCP) agar (bioMérieux; Ref. 43021). Plates were incubated  
112 at 37°C for 18 to 24 h. Cultures were considered pure if only one morphotype was present on the  
113 blood agar plate. For pure cultures, growth on both the CNA and BCP plates was considered as  
114 evidence of the bacteria being gram-positive; growth on only the BCP plate as evidence of the  
115 bacteria being gram-negative. Among gram-negative bacteria, *E.coli* was characterized by a positive  
116 lactose reaction (colour change of the BCP plate from purple to yellow) and a negative urea reaction  
117 (bioMérieux; Ref. 55752), whereas *Klebsiella* was identified by both positive lactose and positive  
118 urea reactions. Gram-positive bacteria were considered to be *Staphylococcus aureus* based on  
119 positive catalase and coagulase reactions (bioMérieux; Ref. 73112) and *Staphylococcus* spp. in the

120 case of positive catalase and negative coagulase results. Identification of *S. uberis* was based on  
121 negative response in the catalase reaction and positive response in the esculin reaction (bioMérieux;  
122 Ref 42086) (National Mastitis Council, 1999). In addition, susceptibility to penicillin was evaluated.  
123 This procedure increases specificity by excluding enterococcal isolates, which are more likely to be  
124 penicillin-resistant than streptococci (Makovec and Ruegg, 2003; Nam *et al.*, 2010). Susceptibility to  
125 penicillin was tested using the disc diffusion method in accordance with the recommendations of the  
126 Société Française de Microbiologie (Soussy, 2013). Using a swab, a Mueller Hinton agar plate  
127 containing 5% sheep blood (MH2, bioMérieux; Ref. 43321) was homogenously plated with a  
128 suspension of *S. uberis* at 0.5 McFarland (equivalent to approximately  $10^8$  colony forming units/ml).  
129 Plates were incubated at 37°C for 18 to 24 h. Bacteria were considered sensitive to penicillin if they  
130 expressed a growth inhibition zone of more than 21 mm around an Oxacillin disc (5 µg; Soussy,  
131 2013). Bacteria that could not be classified using the criteria described here were considered “other  
132 species”.

133

#### 134 ***Cow and Herd Data***

135 Three data sources were used to obtain information about individual cows and their herds of origin,  
136 i.e. (i) private farm records on treatment; (ii) monthly DHI data; and (iii) questionnaires that were  
137 filled out by the farmer and the attending veterinarian. For each cow the date of mastitis diagnosis  
138 (observation of clinical mastitis or notification of SCC data via DHI) and treatment were recorded,  
139 including use of intramammary administration of antimicrobials (IMM), parenteral administration of  
140 antimicrobials (PAR), and use of non-steroidal anti-inflammatory drugs (NSAIDs). DHI data were  
141 collected from the 3 milk recordings preceding the diagnosis of mastitis, the month of diagnosis and  
142 the 3 recordings after diagnosis, if available. This included cow-level SCC data, parity (1 = first  
143 lactation, 2 = second lactation, 3 = third or higher lactation), DIM, milk yield (MY, in kg), fat content  
144 (g/kg), and protein content (g/kg). At herd level, information was collected on the use of pre-milking



145 teat disinfection (PreMTD) and post-milking teat disinfection (PostMTD), use of housing (yes or no)  
146 and, where applicable, on housing type (cubicles or straw yards).

147

148 ***Classification of Cases***

149 Clinical manifestation was classified into four categories based on clinical severity of the current  
150 episode and information on previous episodes of CM in the same animal:

151

- 152 1. Severe: CM with both local and general symptoms ( $T > 39^{\circ}\text{C}$ ; temperature measured on  
153 clinical suspicion of fever by the farmers and results recorded on the data form  
154 accompanying the milk sample);
- 155 2. Non-severe first case ("first"): first episode of CM during the current lactation with local  
156 signs only (abnormalities of milk with or without abnormalities of the udder);
- 157 3. Non-severe repeat case ("repeat"): 2<sup>nd</sup> or subsequent episode of CM during the current  
158 lactation with local signs only in the current episode (abnormalities of milk with or  
159 without abnormalities of the udder);
- 160 4. Subclinical: SCC  $> 200,000$  cells/ml at cow level based on most recent DHI data, not  
161 accompanied by any clinical signs

162

163 For 212 of the 251 *S. uberis* positive animals SCC data were partially available (fewer than 3 records  
164 before or after diagnosis) or complete (3 records before and after diagnosis). SCC data were used to  
165 classify the duration of inflammation prior to diagnosis and the response to treatment after  
166 diagnosis. Inflammation was considered short if at least 2 monthly SCC records prior to diagnosis  
167 were below 200,000 cells/ml (SHORT) and long if at least 2 monthly SCC records exceeded 200,000  
168 cells/ml prior to diagnosis (LONG). An animal was considered cured if at least 2 monthly SCC after  
169 diagnosis were below 200,000 cells/ml (CURE) and not cured if at least 2 monthly SCC exceeded  
170 200,000 cells/ml after diagnosis (NO CURE). For other SCC combinations or missing data, duration

171 and cure were not determined (ND), e.g. for animals in early lactation or for cows that were dried-  
172 off or culled prior to completion of follow-up.

173

#### 174 ***Statistical Analysis***

175 Statistical analyses were performed using Statistix, version 10 (Analytical Software, Tallahassee, FL).

176 Data were inspected for outliers and missing values and descriptive analyses were conducted using

177 tabular and graphical formats. For outcomes of interest with 3 or more categories, data were

178 analysed using categorical methods (Chi-Square analyses), e.g. for cow-level factors associated with

179 clinical severity. The association between clinical severity and milk, fat or protein yield relative to

180 occurrence of mastitis was evaluated using a t-test at each time point. To identify cow- and herd

181 level risk factors for apparent cure as based on SCC, logistic regression was used with backward

182 stepwise analysis. The final logistic regression equation was:

183

184  $\text{Logit (SCC cure)} = \text{intercept} + \text{Clinical manifestation} + \text{Duration} + \text{Parity} + \text{Treatment} + \text{error}$

185

186 where Clinical manifestation is severe, first, repeat or subclinical as defined above, Duration is the

187 inflammation history based on SCC (short, long, ND), parity is parity group (1, 2, 3+), DIM is

188 categorized into early, mid and late lactation (<100, 100-200, 200+) and treatment is treatment for

189 mastitis (IMM, PAR, and NSAIDs, or no treatment). Two way interactions between the main variables

190 were also evaluated for statistical significance. No correction was made for clustering of cases within

191 herd, because the model would not converge when herd was included due to the large number of

192 herds and the limited number of cases per herd. Goodness of fit of the final model was evaluated

193 using the model deviance and the Hosmer-Lemeshow statistic. In the Hosmer-Lemeshow statistic,

194 the data are divided into 10 approximately equal deciles of observed risk. In these deciles the

195 observed and expected number of observation are compared using a Chi-square distribution with

196  $10-2 = 8$  degrees of freedom (Hosmer and Lemeshow, 2013). A low value of the Hosmer-Lemeshow

197 statistic indicates a good fit to the data. In addition, a deviance value that is close to the remaining

198 degrees of freedom implies that there is no evidence of a poor fit of the model to the data.

199

200

**RESULTS**

201

**202 Descriptive Analysis**

203 Of 624 milk samples submitted for culture 251 (40%) were positive for *S. uberis* in pure culture whilst

204 42 samples (7%) were culture negative. The remaining samples tested positive for *Escherichia coli* (n

205 = 108; 17%), *Klebsiella* (n = 12; 2 %), *Staphylococcus aureus* (n = 76; 12%), *Staphylococcus* spp. (n =

206 103; 17%) or other species (n = 32; 5%). Samples positive for *S. uberis* originated from 142 farms.

207 From 99 farms, a single *S. uberis* positive sample was obtained whilst 20 and 23 farms provided 2 or

208 more *S. uberis* positive samples, respectively. All isolates originated from cows with clinical or

209 subclinical mastitis in one quarter with the exception of three cows where *S. uberis* was isolated

210 from two quarters on the same sampling date. The clinical manifestation of *S. uberis* positive

211 mastitis cases was significantly different from the clinical manifestation of *S. uberis* negative cases

212 (Chi-square = 38.0, df = 3,  $P < 0.005$ ; Figure 1), with *S. uberis* overrepresented among non-severe

213 first cases and underrepresented among subclinical cases. Distribution across parities was not

214 different between *S. uberis* and other diagnoses (Chi-square = 1.56, df = 2,  $P = 0.46$ ). During the first

215 100 days of lactation *S. uberis* was less common than other diagnoses, whereas it was more common

216 between 100 and 200 DIM (Chi-square=10.13, df = 2,  $P < 0.05$ ). Milk yield, fat and protein content

217 were not different between *S. uberis* and non *S. uberis* cases prior to infection (results not shown).

218 Cow level data for *S. uberis* cases are summarized in Table 1. Severe and subclinical *S. uberis*

219 cases were overrepresented in parity 1 compared to higher parities, whereas non-severe first cases

220 were overrepresented in parity 2 and repeat cases in higher parities, respectively (Chi-square =

221 13.67, df = 6,  $P < .05$ ). Severe cases were overrepresented in early lactation, whereas repeat cases

222 were overrepresented in mid-lactation (Chi-square = 13.02, df = 6,  $P < 0.05$ ; Table 1). Treatment

223 records were available for ca. 80% of severe, first and repeat cases and for 42% of subclinical cases

224 (Chi-square = 22, df = 3,  $P = 0.0001$ ). When no treatment was recorded, this was considered to

225 indicate that no treatment was administered. Intramammary antibiotics as the only treatment were  
226 more commonly used to treat non-severe first cases as compared to severe, repeated and  
227 subclinical cases, whereas they were more commonly used in combination with parenteral  
228 treatment for repeat cases and subclinical cases (Chi-square = 31; df =6,  $P < 0.0001$ ; Table 1). The  
229 combination of intramammary and parenteral antimicrobials with anti-inflammatory treatment was  
230 mostly used in severe cases and never for subclinical cases (Table 1). Milk production was  
231 numerically lower in severe cases than in non-severe cases, both before and after diagnosis of  
232 clinical or subclinical mastitis, with the exception of yield at 3 DHI recordings prior to diagnosis, but  
233 the difference was not significant. No differences were detected between severity classes with  
234 regard to fat and protein content of milk before or after diagnosis of mastitis (data not shown).

235 Herd-level data was collected on farms with *S. uberis* positive results and is presented in  
236 Table 2. Most herds were housed, either full time or part time. Straw yards were the predominant  
237 housing system, with only 22% of herds housed in cubicles. PostMTD was used in almost all herds  
238 and in more than half of all herds both PreMTD and PostMTD were used. Use of PreMTD without  
239 PostMTD was not reported. Severe cases were overrepresented in herds without PostMTD (Chi-  
240 square = 10.23, df = 3,  $P < .05$ ).

241

#### 242 ***Factors Associated with Cure of S. uberis IMI***

243 Cure was evaluated based on post diagnosis SCC values and results from the regression model are  
244 shown in Table 3. A total of 125 cases had complete data and were included in this analysis. Model  
245 deviance was 127.4 on 115 degrees of freedom, i.e. the values were similar and there was no  
246 indication of a poor fit of the model to the data. In addition, the Hosmer-Lemeshow statistic was low  
247 (6.05), implying a good fit to the data. The probability of cure was significantly higher in animals in  
248 lactation 1 and 2 compared to older animals. The probability of cure increased numerically with  
249 increasing number of treatment types, i.e. from no treatment to intramammary antimicrobials only

250 to combined intramammary and parenteral antimicrobials, to both routes of antimicrobial  
251 administration combined with NSAID. However, there was no statistically significant difference in  
252 cure between treatments. Finally, the probability of cure was higher among IMI with a short history  
253 of inflammation than those with a long history of inflammation prior to treatment (Table 3). Clinical  
254 manifestation and herd level variables were not associated with cure.

255

256

**DISCUSSION**

257 In this study we aimed to use routinely available herd and animal-level data to support control of *S.*  
258 *uberis* mastitis and the judicious use of antimicrobials. Risk factors for the incidence of *S. uberis*  
259 mastitis (clinical mastitis or IMI) have been described at herd-level (Barkema *et al.*, 1999; Ericsson  
260 Unnerstad *et al.*, 2009) and animal-level (Zadoks *et al.*, 2001; Breen *et al.*, 2009), and the impact of  
261 different treatment regimens on the outcome of treatment of *S. uberis* mastitis has been described  
262 for experimentally induced (Hillerton and Kliem, 2002; Oliver *et al.*, 2003) and naturally occurring  
263 infections (reviewed in Zadoks, 2007). To our knowledge, animal-level risk factors for severity of  
264 disease or treatment outcome of *S. uberis* IMI have not been described. Here, we show for the first  
265 time that animal-level data can be used to predict the outcome of antimicrobial treatment of *S.*  
266 *uberis* mastitis and to guide treatment decisions. Specifically, the probability of cure was higher  
267 among 1st and 2nd parity animals compared to older cows, and in animals with at most a single  
268 elevated cow-level SCC before diagnosis compared to those with multiple high SCC records. Those  
269 findings are strikingly similar to results obtained for *Staphylococcus aureus* IMI across a range of  
270 studies covering both clinical and subclinical mastitis (reviewed in Barkema *et al.*, 2006) and can be  
271 used to inform decisions about treatment duration or the choice between treatment and culling. The  
272 individual making treatment decisions will be able to weigh these factors in the decision making and  
273 to use this information to provide a realistic prognosis. Pathogen-specific predictors for cure, as  
274 described here for *S. uberis*, are particularly useful when information on the causative agent is  
275 available. Several studies have demonstrated the feasibility of using on-farm diagnostics to inform  
276 case management (Lago *et al.*, 2011; Cameron *et al.*, 2014) and additional tests for rapid or on-farm  
277 screening of milk samples are under development, including culture and DNA-based tests (Viora *et*  
278 *al.*, 2014; Bosward *et al.*, 2016). Considering the similarities between results obtained for *S. uberis*  
279 and *S. aureus*, some of this information may also be of value in the absence of an etiological  
280 diagnosis, although further field evaluation will be needed to validate such a generic approach.

281 Increased parity was associated with a reduced likelihood of cure. This is not merely a  
282 reflection of the chronicity of infection, because parity was significant after correction for SCC, which  
283 is a proxy for duration (Barkema et al., 2006). The mechanism behind reduced probability of cure in  
284 older animals is unknown. Possible explanations were discussed by Barkema and co-workers (2006)  
285 for the response to treatment of *Staph. aureus* IMI. One potential explanation is the change in ratio  
286 between udder volume, which increases with age, and the administered dose of antimicrobials,  
287 which is independent of age, resulting in a lower dose per unit udder volume in older animals  
288 (Barkema et al., 2006). This reasoning would also apply to *S. uberis* treatment. Immunosenescence,  
289 the waning of the immune response with age, could be postulated to play a role in deterioration of  
290 treatment outcome with age, but there is no specific evidence for this in the context of bovine  
291 mastitis. Regardless of the underlying mechanism, the comparatively poor treatment response of  
292 older cows can be interpreted as an imperative to help our cows to age healthily, e.g. by selecting  
293 for cows with high genetic merit for udder health or immune responsiveness (Thompson-Crispi et al.,  
294 2014). In addition, animal-level risk factors should be minimized where possible. For example,  
295 severe teat end hyperkeratosis is an animal-level risk factor for *S. uberis* CM, and the risk of  
296 hyperkeratosis can be reduced by avoiding overmilking (Breen et al., 2009; Edwards et al., 2013).

297 The observation that duration of IMI, as measured by number of elevated monthly SCC prior  
298 to treatment, is predictive of cure is compatible with previous data on both *S. uberis* and *Staph.*  
299 *aureus*. A detailed longitudinal study of *S. uberis* IMI in 2 herds showed that some episodes of CM  
300 are due to recent IMI, whereas other CM episodes are preceded by periods of elevated (Zadoks et  
301 al., 2003). CM episodes without preceding SCC elevation were more likely to be followed by cure  
302 than CM episodes with preceding SCC elevation. Similarly, in several treatment trials of *Staph.*  
303 *aureus* IMI, higher or longer SCC elevation prior to treatment was associated with a decreased  
304 probability of cure (reviewed in Barkema et al., 2006). A poor response of chronic *Staph. aureus* IMI  
305 to treatment may be explained in part by micro-abscess formation and fibrosis (Erskine et al., 2003).  
306 Fibrosis also occurs during *S. uberis* mastitis, starting as early as 6 days after infection in



307 experimental challenge studies. It is accompanied by presence of the pathogen in subepithelial and  
308 septal tissue and lymphatic vessels and lymph nodes (Thomas *et al.*, 1994). This may explain why the  
309 response of *S. uberis* mastitis to treatment can be poor, even after extended therapy (Milne *et al.*,  
310 2005). Both in experimentally induced and in persistent *S. uberis* IMI, extended therapy increases  
311 the probability of cure (Oliver *et al.*, 2003; Swinkels *et al.*, 2014). The benefits of extended therapy  
312 must be weighed against its disadvantages, including increased costs of antibiotics and milk discard,  
313 and increased risk of residue in milk and selection for antimicrobial resistance (Hillerton and Kliem,  
314 2002; Barkema *et al.*, 2006). As in any risk factor study, the risk factors identified in the current  
315 study, including treatment modality, and their coefficients allow us to quantify the increase or  
316 decrease in the likelihood of a particular treatment outcome, but the specific outcome in any  
317 individual animal cannot be predicted.

318           In the current study a numerical but non-significant increase in cure was observed with an  
319 increase in treatment modalities (intramammary and parenteral antimicrobials and NSAIDs). This  
320 study was not, however, a randomized controlled clinical trial, nor was it meant to be. Farmers  
321 tended to treat severe cases of mastitis with a combination of intramammary, parenteral and anti-  
322 inflammatory products, first clinical cases with intra-mammary treatment only and repeated- and  
323 subclinical cases with antimicrobial treatment by both the intra-mammary and parenteral route.  
324 This implies farmers' awareness of the usefulness of cow-specific treatment, with consideration of  
325 both animal welfare and economic aspects of treatment. This information provides evidence that  
326 farmers are willing to make cow-specific decisions and bodes well for the feasibility of including cow-  
327 specific risk factors in future protocols. In our practice, farm specific treatment protocols are already  
328 discussed with each farmer on an annual basis and the treatment choices reported by the farmers  
329 are in line with those protocols. As a next step towards judicious use of antimicrobials, we envisage  
330 implementation of cow-specific protocols.

331           Animals with short duration mastitis (no or single SCC elevation prior to diagnosis) were  
332 likely to cure (no or single SCC elevation after diagnosis), whereas animals with long duration  
333 mastitis (multiple SCC elevation prior to diagnosis) were likely not to cure (multiple SCC elevations  
334 after diagnosis). Similarly, data availability post-diagnosis mirrored data availability pre-diagnosis, i.e.  
335 animals with incomplete SCC data prior to diagnosis often had incomplete SCC data after diagnosis  
336 too (data not shown). This would mostly apply to animals in early lactation that were lost to follow-  
337 up due to culling. Thus, although our analysis shows no significant difference in cure between  
338 different severity classes, this result is affected by “healthy worker bias”, whereby only surviving  
339 cows are included in the analysis. Indeed, loss to follow-up as indicated by absence of data on cure  
340 was proportionally higher for severe cases than for non-severe cases (Table 1).

341           Of the herd-level factors considered in this study, use of PostMTD was associated with a  
342 reduced risk of severe, repeat and subclinical *S. uberis* mastitis compared to first cases of mastitis.  
343 The value of PostMTD in reducing the risk of *S. uberis* IMI has been documented repeatedly (Zadoks  
344 et al., 2003; Galton, 2004; Williamson and Lacy-Hulbert, 2013) but it has not been linked to clinical  
345 manifestation. Strain-specific transmission and virulence patterns have previously been suggested or  
346 documented (Zadoks et al., 2003; Tassi et al., 2013) and could theoretically contribute to an  
347 association between PostMTD and clinical manifestation. It has also been hypothesized that host  
348 immune status may contribute to clinical manifestation of *S. uberis* IMI (Tassi et al., 2013). Indeed, in  
349 the current study, severe cases of CM were overrepresented among heifers and animals in early  
350 lactation. This emphasizes the importance of another herd-level management factor, i.e. adequate  
351 care for non-lactating animals. Considering that *S. uberis* is common in the faeces and environment  
352 of cattle (Zadoks et al., 2005), environmental hygiene is of particular importance. The risk of  
353 infection in heifers and dry cows can also be reduced through use of teat spray and internal teat  
354 sealants, respectively (Lopez-Benavides et al., 2009; Compton et al., 2014). With increasing pressure  
355 to reduce antimicrobial use, implementation of non-antimicrobial mastitis prevention measures  
356 becomes increasingly important.

357 In this field study, definitions of transient and persistent IMI and cure were based on SCC  
358 data. Although repeated post-treatment culture has been considered the “gold standard” for cure in  
359 clinical trials, additional or alternative metrics for cure are increasingly reported in field studies. SCC  
360 has been used as a primary criterion for cure in studies of clinical mastitis, subclinical mastitis, and  
361 dry cow treatment (St. Rose *et al.*, 2003; Lago *et al.*, 2011; Persson *et al.*, 2015). SCC is routinely used  
362 as an indicator of infection status (Schukken *et al.*, 2003), although the probability of bacteriological  
363 cure is higher than the probability of SCC-based cure in studies of chronic streptococcal mastitis (St.  
364 Rose *et al.*, 2003). SCC is of immediate interest to farmers, unlike bacteriological cure which is  
365 primarily of academic interest. Moreover, SCC is routinely available at very low cost, which makes its  
366 large-scale use feasible in field studies, veterinary practice and farm management. Finally, SCC  
367 captures long-term outcomes of mastitis treatment, whereas culture results generally only reflect  
368 the first few weeks post-treatment. Thus, SCC is a convenient, affordable and meaningful indicator  
369 of treatment outcome.

370 In conclusion, we show that treatment recommendations can be informed by animal-level  
371 data that is routinely available to farmers and veterinarians, such as parity and SCC. To some extent,  
372 treatment recommendations can be animal-specific rather than pathogen-specific, as both *S. uberis*  
373 IMI and *Staph. aureus* IMI show a better response to treatment in animals in first or second lactation  
374 and in animals with a single high SCC than in older animals or animals with multiple high SCC values  
375 prior to treatment. In older animals or animals with multiple high SCC values the simultaneous use of  
376 multiple treatment modalities may enhance the probability of cure but this would result in increased  
377 use of antimicrobials. To limit the need for such treatment, continued or renewed emphasis on herd  
378 management and infection prevention is needed. Formal validation of the observations described  
379 here through a randomized controlled clinical trial may strengthen the evidence base underpinning  
380 the suggested treatment decisions. In the absence of such validation, the evidence presented here is  
381 the best available evidence to inform decisions on treatment of *S. uberis* mastitis, the most common  
382 type of mastitis observed in this and many other studies.

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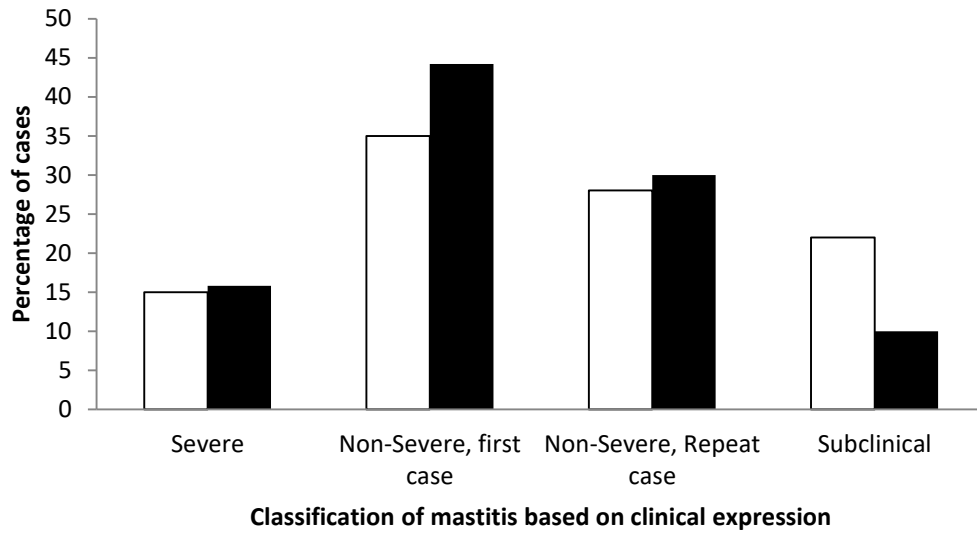


497 **Figure legend**

498 **Figure 1.** Clinical manifestation of mastitis for quarters with *S. uberis* negative (n = 373, white) and *S.*  
499 *uberis* positive (n =251, black) milk samples (Chi-square = 38.0, df = 3,  $P < 0.005$ ). Severe: clinical  
500 mastitis (CM) with local and general symptoms ( $T > 39^{\circ}\text{C}$ ; checked upon clinical suspicion of fever);  
501 Non-severe first case: first occurrence of CM during the current lactation with local signs only  
502 (abnormalities of milk with or without abnormalities of the udder). Non-severe repeat case: repeat  
503 occurrence of CM during the current lactation with local signs only during the current episode;  
504 Subclinical\_mastitis: elevated cow-level SCC ( $> 200.000$  cells/ml based on DHI data) not accompanied  
505 by any visible abnormalities.

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Samson *et al.*, Figure 1.

523 **Table 1.** Cow- level data for *S. uberis* positive mastitis cases (number and (%)) with break-down by  
 524 manifestation. Severe: clinical mastitis (CM) with local and general symptoms (T > 39°C; checked  
 525 upon clinical suspicion of fever); First: first occurrence of CM during the current lactation with local  
 526 signs only (abnormalities of milk with or without abnormalities of the udder). Repeat: repeat  
 527 occurrence of CM during the current lactation with local signs only during current episode;  
 528 Subclinical: cow-level SCC (> 200,000 cells/ml based on DHI data) not accompanied by any signs.

529

Cow factor	All <i>S. uberis</i> cases, n (%)	<i>S. uberis</i> cases by clinical manifestation, n (%)			
		Severe	First	Repeat	Subclinical
DIM					
<100	123 (100)	26 (21)	56 (46)	27 (22)	14 (11)
100 to 200	63 (100)	7 (11)	25 (40)	27 (43)	4 (6)
> 200	50 (100)	4 (8)	24 (48)	17 (34)	5 (10)
<i>subtotal</i>	236 (100)	37 (16)	105 (44)	71 (30)	23 (10)
Parity					
First	61 (100)	12 (20)	23 (38)	16 (26)	10 (16)
Second	55 (100)	7 (13)	34 (62)	12 (22)	2 (4)
Higher	122 (100)	17 (14)	49 (40)	44 (36)	12 (10)
<i>subtotal</i>	238 (100)	36 (15)	106 (45)	72 (30)	24 (10)
Treatment <sup>1</sup>					
none	49 (100)	7 (14)	17 (35)	12 (24)	13 (27)
IMM	67 (100)	6 (9)	42 (63)	17 (25)	2 (3)
IMM + PAR	103 (100)	14 (14)	41 (40)	40 (39)	8 (12)
IMM + PAR + NSAID	19 (100)	10 (53)	6 (32)	3 (16)	0 (0)
<i>subtotal</i>	238 (100)	37 (16)	106 (45)	72 (30)	23 (10)

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Duration<sup>2</sup>

Short	78 (100)	8 (10)	35 (45)	30 (38)	5 (6)
Long	66 (100)	8 (12)	25 (38)	23 (35)	10 (15)
ND	96 (100)	22 (23)	46 (48)	19 (20)	9 (9)
<i>subtotal</i>	240 (100)	38 (16)	106 (44)	72 (30)	24 (10)

Cure

Yes	79 (100)	10 (13)	39 (49)	23 (29)	7 (9)
No	97 (100)	12 (12)	38 (39)	33 (34)	14 (14)
ND	64 (100)	16 (25)	29 (45)	16 (25)	3 (5)
<i>subtotal</i>	240 (100)	38 (16)	106 (44)	72 (30)	24 (10)

530

531 <sup>1.</sup> Treatment: IMM = intra-mammary antibiotic treatment, PAR= parenteral antibiotic treatment,

532 NSAID = non-steroidal anti-inflammatory drugs.

533 <sup>2.</sup> Duration: Short = at least 2 monthly cow-level SCC < 200,000 cells/ml before diagnosis; Long =

534 at least 2 monthly cow-level SCC > 200,000 cells/ml before diagnosis; ND = not determined due

535 to insufficient SCC data before diagnosis.

536 <sup>3.</sup> Cure: Yes = at least 2 monthly cow-level SCC < 200,000 cells/ml after diagnosis; Long = at least 2

537 monthly cow-level SCC > 200,000 cells/ml after diagnosis; ND = not determined due to

538 insufficient SCC data after diagnosis.

539 **Table 2.** Herd-level data for *S. uberis* positive mastitis cases (number and (%)) with breakdown by  
 540 manifestation. Severe: clinical mastitis (CM) with local and general symptoms (T > 39°C; checked  
 541 upon clinical suspicion of fever); First: first occurrence of CM during the current lactation with local  
 542 signs only (abnormalities of milk with or without abnormalities of the udder). Repeat: repeat  
 543 occurrence of CM during the current lactation with local signs only during current episode;  
 544 Subclinical: cow-level SCC (> 200.000 cells/ml based on DHI data) not accompanied by any signs.

545

Herd factor	All <i>S. uberis</i> cases, n (%)	<i>S. uberis</i> cases by clinical manifestation, n (%)			
		Severe	First	Repeat	Subclinical
Housing					
Permanent	120 (100)	15 (13)	57 (48)	39 (33)	9 (8)
Partial	92 (100)	20 (22)	32 (35)	28 (30)	12 (13)
None	28 (100)	3 (29)	17 (61)	5 (18)	3 (11)
<i>subtotal</i>	240 (100)	38 (16)	106 (44)	72 (30)	24 (10)
Bedding					
Straw yard	187 (100)	27 (11)	84 (45)	58 (31)	18 (75)
Cubicles	53 (100)	11 (21)	22 (42)	14 (26)	6 (25)
<i>Subtotal</i>	240 (100)	38 (16)	106 (44)	72 (30)	24 (100)
Pre-dipping					
Yes	130 (100)	19 (15)	56 (43)	44 (34)	11 (8)
No	106 (100)	18 (17)	48 (45)	27 (25)	13 (12)
<i>Subtotal</i>	236 (100)	37 (16)	104 (44)	71 (30)	24 (10)
Post-dipping					
Yes	214 (100)	29 (14)	94 (44)	67 (31)	24 (11)
No	22 (100)	8 (36)	10 (45)	4 (18)	0 (0)
<i>Subtotal</i>	236 (100)	37 (16)	104 (44)	71 (30)	24 (10)

546

547 **Table 3.** Logistic regression of cow-factors versus cure for 125 cases of *S. uberis* mastitis.

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Variable	Coefficient (SE)	Odds Ratio	95% C.I. for		
			Odds Ratio	Z-value	P-value
Constant	-4.8 (1.3)			-3.68	0.0002
Parity					
First	1.7 (0.6)	5.5	1.8 to 5.5	3.1	0.0023
Second	1.3 (0.6)	3.8	1.3 to 11.1	2.42	0.016
Third or higher	Base				
Clinical manifestation <sup>1</sup>					
Severe	1.6 (1.3)	4.8	0.3 to 68.7	1.16	0.24
First	2.0 (1.2)	7.1	0.7 to 77.0	1.65	0.10
Repeat	1.9 (1.2)	6.4	0.6 to 70.9	1.54	0.13
Subclinical	Base				
Duration <sup>2</sup>					
Short	1.7 (0.5)	3.1	2.2 to 14.2	3.74	0.0002
Long	Base				
Treatment <sup>3</sup>					
IMM + PAR + NSAID	2.2 (1.3)	9.4	0.7 to 9.4	1.68	0.09
IMM + PAR	1.1 (0.6)	3.1	0.9 to 3.1	1.79	0.074
IMM	0.7(0.7)	2.1	0.6 to 8.2	1.09	.28
None	Base				

549

550 <sup>1.</sup> Severe: clinical mastitis (CM) with local and general symptoms (T > 39°C; checked upon clinical  
 551 suspicion of fever); First: first occurrence of CM during the current lactation with local signs only  
 552 (abnormalities of milk with or without abnormalities of the udder). Repeat: repeat occurrence

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553 of CM during the current lactation with local signs only during current episode; Subclinical: cow-  
554 level SCC (> 200,000 cells/ml based on DHI data) not accompanied by any signs.

555 <sup>2.</sup> Duration: Short = at least 2 monthly cow-level SCC < 200,000 cells/ml before diagnosis; Long =  
556 at least 2 monthly cow-level SCC > 200,000 cells/ml before diagnosis; ND = not determined due  
557 to insufficient SCC data before diagnosis.

558 <sup>3.</sup> Treatment: IMM = intra-mammary antibiotic treatment, PAR= parenteral antibiotic treatment,  
559 NSAID = non-steroidal anti-inflammatory drugs.